CRF Errors Corrected by the STIC Systems Branch CRF Processing Date: Serlai Number: Edited by: Verifled by: _ Changed a file from non-ASCII to ASCII Changed the margins in cases where the sequence text was "wrapped own to the next line. Edited a format error in the Current Application Data section, specifically: Edited the Current Application Data section with the actual current number. The number inputted by the applicant was
the prior application data; or other ____ Added the mandatory heading and subheadings for "Current Application Data". Edited the "Number of Sequences" field. The applicant spelled out a number instead of using an integer. Changed the spelling of a mandatory field (the headings or subheadings), specifically: Corrected the SEQ ID NO when obviously incorrect. The sequence numbers that were edited were: Inserted of corrected a nucleic number at the end of a nucleic line. SEQ ID NO's edited: Corrected subheading placement. All responses must be on the same line as each subheading. If the applicant placed a response below the subheading, this was moved to its appropriate place. Inserted colons after headings/subheadings. Headings edited included: Deleted extra, invalid, headings used by an applicant, specifically: Deleted: non-ASCII "garbage" at the beginning/end of files; secretary initials/filename at end of file; page numbers throughout text; other invalid text, such as___ Inserted mandatory headings, specifically: Corrected an obvious error in the response, specifically: Edited identifiers where upper case is used but lower case is required, or vice versa. Corrected an error in the Number of Sequences field, specifically: A "Hard Page Break" code was inserted by the applicant. All occurrences had to be deleted. Deleted ending stop codon in amino acid sequences and adjusted the "(A)Length:" field accordingly (error due to a Patentin bug). Sequences corrected: _ Other:

*Examiner: The above corrections must be communicated to the applicant in the first Office Action. DO NOT send a copy of this form.



OIPE

RAW SEQUENCE LISTING

PATENT APPLICATION: US/09/904,992

DATE: 02/17/2002 TIME: 14:31:34

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         Wood, William, I.
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VERIFICATION SUMMARY

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DATE: 02/08/2002

PATENT APPLICATION: US/09/904,992

3 <110> APPLICANT: Genentech, Inc.

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- 80 <151> PRIOR FILING DATE: 1999-12-20
- 82 <150> PRIOR APPLICATION NUMBER: PCT/US99/30999
- 83 <151> PRIOR FILING DATE: 1999-12-20
- 84 <150> PRIOR APPLICATION NUMBER: PCT/US00/00219
- 85 <151> PRIOR FILING DATE: 2000-01-05
- 87 <160> NUMBER OF SEQ ID NOS: 423

ERRORED SEQUENCES

- 5293 <210> SEQ ID NO: 173
- 5294 <211> LENGTH: 43
- 5295 <212> TYPE: DNA
- 5296 <213> ORGANISM: Artificial Sequence
- 5298 <220> FEATURE:
- 5299 <223> OTHER INFORMATION: Description of Artificial Sequence: Synthetic
- 5300 oligonucleotide probe
- 5302 <400> SEQUENCE: 173
- E--> 5303 ggactcactg gcccaggcct tcaatatcac cagccaggac gat

VERIFICATION SUMMARY

L:5479~M:341~W: (46) "n" or "Xaa" used, for SEQ ID#:175 L:6540 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:206

DATE: 02/08/2002 TIME: 14:55:56

PATENT APPLICATION: US/09/904,992

Input Set : N:\EBONY'S\EP.txt

Output Set: N:\CRF3\02082002\1904992.raw

L:34 M:270 C: Current Application Number differs, Replaced Current Application Number L:35 M:271 C: Current Filing Date differs, Replaced Current Filing Date L:511 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:13 L:512 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:13 L:513 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:13 L:514 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:13 $L:769\ M:341\ W:$ (46) "n" or "Xaa" used, for SEQ ID#:26 L:1701 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:50 L:3586 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:113 L:4040 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:131 L:5303 M:254 E: No. of Bases conflict, LENGTH:Input:42 Counted:43 SEQ:173 L:5344 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:174